SEQUENCE LISTING

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XU, JUN

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Gly Glu Arg Ser Arg Lys Arg Arg Ala Val Leu Thr Gln Lys Gln Lys
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Gly Arg Gly Leu Gln Ala Gln Gly Tyr Gly Val Arg Ile Gln Asp Ala
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Gly Val Tyr Leu Leu Tyr Ser Gln Val Leu Phe Gln Asp Val Thr Phe
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Thr Met Gly Gln Val Val Ser Arg Glu Gly Gln Gly Arg Gln Glu Thr
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Leu Phe Arg Cys Ile Arg Ser Met Pro Ser His Pro Asp Arg Ala Tyr
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Asn Ser Cys Tyr Ser Ala Gly Val Phe His Leu His Gln Gly Asp Ile
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Glu Ser Gln Phe Gln Arg His Met Arg Asp His Glu Arg Asn Asp Lys
     50                  55                  60

Pro His Arg Cys Asp Gln Cys Pro Gln Thr Phe Asn Val Glu Phe Asn
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Leu Thr Leu His Lys Cys Thr His Ser Gly Glu Asp Pro Thr Cys Pro
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Gln Glu Leu Ala Gly Thr Arg Gln His Ala Cys Lys Ala Cys Lys Lys
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Glu Phe Glu Thr Ser Ser Glu Leu Lys Glu His Met Lys Thr His Tyr
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Lys Ile Arg Val Ser Ser Thr Arg Ser Tyr Asn Arg Asn Ile Asp Arg
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Ser Gly Phe Thr Tyr Ser Cys Pro His Cys Gly Lys Thr Phe Gln Lys
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Pro Ser Gln Leu Thr Arg His Ile Arg Ile His Thr Gly Glu Arg Pro
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Pro Val Glu Ser Gly Gln Ser Pro Gln Pro Gly Gln Gln Leu Ser Ile
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gggctgtctt caattccagc tgcagcacat cctaatgact cctgccatgc caagacctct 1260
gcaccacacg ctcaaaaccc agatgtttcc agcgtttcaa atgagcagac ggaccccaca 1320
gacgcagagc aagaaaaaga acaggaaagc ccggagaaac tggataaaaa aaaaaaaaaa 1380
agggccacat gtgctcgagc tgcaggtcgc ggccgctag                        1419
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<210> 9
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
oligonucleotide

<400> 9
aacccctgcc acaacggtgg t 21

<210> 10
<211> 21
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 10
aaccccugcc acaacggugg u 21

<210> 11
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 11
aaccactgtg agacgaaatg t 21

<210> 12
<211> 21
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 12
aaccacugug agacgaaaug u 21

<210> 13
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 13
aactgcccca gcgatctctg c 21

<210> 14
<211> 21
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 14
aacugcccca gcgaucucug c 21

<210> 15
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 15
aacctaattc tcctgaggct g 21

<210> 16
<211> 21
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 16
aaccuaauuc uccugaggcu g 21

<210> 17
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 17
aatgcggaga acactaatta t 21

<210> 18
<211> 21
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 18
aaugcggaga acacuaauua u 21

<210> 19
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 19
aatgacaagc cacatcgatg t 21

<210> 20
<211> 21
<212> RNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 20
aaugacaagc cacaucgaug u 21

<210> 21
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 21
aagctggaca ttccctctgc g 21

<210> 22
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 22
aagagcccag cttcctgcag c 21

<210> 23
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 23
aactgttgag gagcccatgg a 21

<210> 24
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 24
aatctgatga tgaagctgca g 21

<210> 25
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 25
aagagcccag cttcctgcag c 21

<210> 26
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 26
aagctggaca ttccctctgc g 21

<210> 27
<211> 152
<212> PRT
<213> Homo sapiens

<400> 27
Ala Leu Arg Asn Trp Gln Val Tyr Arg Leu Val Thr Tyr Ile Phe Val
1 5 10 15

Tyr Glu Asn Pro Ile Ser Leu Leu Cys Gly Ala Ile Ile Ile Trp Arg
20 25 30

Phe Ala Gly Asn Phe Glu Arg Thr Val Gly Thr Val Arg His Cys Phe
35 40 45

Phe Thr Val Ile Phe Ala Ile Phe Ser Ala Ile Ile Phe Leu Ser Phe
50 55 60

Glu Ala Val Ser Ser Leu Ser Lys Leu Gly Glu Val Glu Asp Ala Arg
65 70 75 80

Gly Phe Thr Pro Val Ala Phe Ala Met Leu Gly Val Thr Thr Val Arg
85 90 95

Ser Arg Met Arg Arg Ala Leu Val Phe Gly Met Val Val Pro Ser Val
100 105 110

Leu Val Pro Trp Leu Leu Leu Gly Ala Ser Trp Leu Ile Pro Gln Thr
115 120 125

Ser Phe Leu Ser Asn Val Cys Gly Leu Ser Ile Gly Leu Ala Tyr Ala
130 135 140

His Leu Leu Leu Phe His Arg Pro
145 150

<210> 28
<211> 152
<212> PRT
<213> Saccharomyces cerevisiae

<400> 28
Leu Leu Gln Lys Arg Gln Leu Tyr Glu Ile Ile Thr Tyr Val Thr Leu
1 5 10 15

His Leu Ser Met Leu His Ile Val Phe Asn Phe Val Ser Leu Leu Pro
20 25 30

Ala Met Ser Gln Phe Glu Lys Lys Gln Gly Thr Leu Ala Cys Ile Leu
35 40 45

Val Thr Val Ile Pro Tyr Thr Leu Phe Pro Gly Ile Met His Leu Ile
50 55 60

Val Tyr His Phe Phe Leu Arg Lys Asp Tyr Val Ser Ile Ala Gly Leu
65 70 75 80

Ser Gly Trp Ala Phe Ala Phe Ile Ser Ala Ser Cys Val His Ser Pro
85 90 95

```
Gln Arg Leu Ile Ser Phe Phe Asn Leu Phe Ser Ile Pro Ala Tyr Cys
            100                 105                 110

Phe Pro Ile Ile Tyr Leu Ile Met Thr Thr Ile Leu Val Pro Lys Ala
        115                 120                 125

Ser Phe Ile Gly His Ala Ser Gly Ala Val Met Gly Tyr Cys Thr Pro
    130                 135                 140

Phe Met Leu Gly Ser Ile Pro Leu
145                 150
```

<210> 29
<211> 145
<212> PRT
<213> Schizosaccharomyces pombe

<400> 29

```
Pro Arg Ser Leu Glu Gly Leu Arg Gly Ile Val Phe Ala Pro Phe Leu
  1               5                  10                  15

His Ala Asp Phe Gly His Leu Ile Ala Asn Ser Val Pro Phe Val Val
             20                  25                  30

Leu Ala Trp Leu Val Met Leu Gln Glu Val Ser Asp Phe Trp Ile Val
         35                  40                  45

Thr Ile Ile Thr Met Val Val Gly Gly Leu Gly Val Trp Leu Ile Ala
     50                  55                  60

Pro Pro Asn Thr Val Thr Val Gly Ala Ser Ile Leu Ile Phe Gly Tyr
 65                  70                  75                  80

Leu Gly Phe Leu Leu Phe Arg Gly Trp Phe Gln Lys Asn Leu Ala Ser
                 85                  90                  95

Ile Val Leu Ser Ile Val Val Leu Val Leu Tyr Gly Ser Ala Leu Trp
            100                 105                 110

Gly Leu Leu Pro Gly Arg Ala Gly Val Ser Trp Gln Gly His Leu Phe
        115                 120                 125

Gly Phe Ile Gly Gly Ala Ile Ala Ala Trp Leu Ile Ala Arg Glu Lys
    130                 135                 140

His
145
```

<210> 30
<211> 145
<212> PRT
<213> Saccharomyces cerevisiae

<400> 30

```
Ser Lys Ser Asn Ala Arg Pro Val Val Ala Ile Gly Asp Ser Asp Ile
  1               5                  10                  15
```

```
Tyr Ser Tyr Arg Leu Trp Ser Phe Phe Cys Gln Trp Ile Asn Thr Ile
            20                  25                  30

Phe Cys Trp Ser Asn Arg Arg Arg Pro Leu Gly Leu Thr Pro Phe Leu
        35                  40                  45

Leu Leu Tyr Val Leu Ser Gly Val Met Gly Asn Ala Phe Thr Phe Trp
    50                  55                  60

Leu Thr Pro Glu Thr Val Ala Ala Gly Ala Ser Thr Ser Leu Phe Gly
65                  70                  75                  80

Leu Phe Ala Ala Ile Val Val Leu Ser Phe Leu Gly Lys Asn Gln Ala
                85                  90                  95

Leu Lys Asp Leu Gly Lys Ser Tyr Gln Thr Leu Ile Val Val Asn Leu
            100                 105                 110

Leu Met Asn Leu Phe Met Pro Asn Val Ser Met Ala Gly His Ile Gly
        115                 120                 125

Gly Val Val Gly Gly Ala Leu Leu Ser Ile Val Phe Pro Thr Lys Met
    130                 135                 140

Arg
145
```

<210> 31
<211> 156
<212> PRT
<213> Homo sapiens

<400> 31

```
Pro Glu Lys Arg Glu Glu Ala Trp Arg Phe Ile Ser Tyr Met Leu Val
  1               5                  10                  15

His Ala Gly Val Gln His Ile Leu Gly Asn Leu Cys Met Gln Leu Val
            20                  25                  30

Leu Gly Ile Pro Leu Glu Met Val His Lys Gly Leu Arg Val Gly Leu
        35                  40                  45

Val Tyr Leu Ala Gly Val Ile Ala Gly Ser Leu Ala Ser Ser Ile Phe
    50                  55                  60

Asp Pro Leu Arg Tyr Leu Val Gly Ala Ser Gly Gly Val Tyr Ala Leu
65                  70                  75                  80

Met Gly Gly Tyr Phe Met Asn Val Leu Val Asn Phe Gln Glu Met Ile
                85                  90                  95

Pro Ala Phe Gly Ile Phe Arg Leu Leu Ile Ile Ile Leu Ile Ile Val
            100                 105                 110

Leu Asp Met Gly Phe Ala Leu Tyr Arg Arg Phe Phe Val Pro Glu Asp
        115                 120                 125
```

Gly Ser Pro Val Ser Phe Ala Ala His Ile Ala Gly Gly Phe Ala Gly
130 135 140

Met Ser Ile Gly Tyr Thr Val Phe Ser Cys Phe Asp
145 150 155

<210> 32
<211> 145
<212> PRT
<213> Escherichia coli

<400> 32
Pro Thr Leu Lys Phe Glu Phe Trp Arg Tyr Phe Thr His Ala Leu Met
1 5 10 15

His Phe Ser Leu Met His Ile Leu Phe Asn Leu Leu Trp Trp Trp Tyr
20 25 30

Leu Gly Gly Ala Val Glu Lys Arg Leu Gly Ser Gly Lys Leu Ile Val
35 40 45

Ile Arg Ser Ile Ser Ala Leu Leu Ser Gly Tyr Val Gln Gln Lys Phe
50 55 60

Ser Gly Pro Trp Phe Gly Gly Leu Ser Gly Val Val Tyr Ala Leu Met
65 70 75 80

Gly Tyr Val Trp Leu Arg Gly Glu Arg Asp Pro Gln Ser Gly Ile Tyr
85 90 95

Leu Gln Arg Gly Leu Ile Ile Phe Ala Leu Ile Trp Ile Val Ala Gly
100 105 110

Trp Phe Asp Leu Phe Gly Met Ser Met Ala Asn Gly Ala His Ile Ala
115 120 125

Gly Leu Ala Val Gly Leu Ala Met Ala Phe Val Asp Ser Leu Asn Ala
130 135 140

Arg
145

<210> 33
<211> 157
<212> PRT
<213> Homo sapiens

<400> 33
Ser Asn Pro Ala Ser Lys Val Leu Cys Ser Pro Met Leu Leu Ser Thr
1 5 10 15

Phe Ser His Phe Ser Leu Phe His Met Ala Ala Asn Met Tyr Val Leu
20 25 30

```
Trp Ser Phe Ser Ser Ser Ile Val Asn Ile Leu Gly Gln Glu Gln Phe
          35                  40                  45

Met Ala Val Tyr Leu Ser Ala Gly Val Ile Ser Asn Phe Val Ser Tyr
      50                  55                  60

Leu Gly Lys Val Ala Thr Gly Arg Tyr Gly Pro Ser Leu Gly Ala Ser
 65                  70                  75                  80

Gly Ala Ile Met Thr Val Leu Ala Ala Val Cys Thr Lys Ile Pro Glu
                 85                  90                  95

Gly Arg Leu Ala Ile Ile Phe Leu Pro Met Phe Thr Phe Thr Ala Gly
            100                 105                 110

Asn Ala Leu Lys Ala Ile Ile Ala Met Asp Thr Ala Gly Met Ile Leu
        115                 120                 125

Gly Trp Lys Phe Phe Asp His Ala Ala His Leu Gly Gly Ala Leu Phe
    130                 135                 140

Gly Ile Trp Tyr Val Thr Tyr Gly His Glu Leu Ile Trp
145                 150                 155


<210> 34
<211> 142
<212> PRT
<213> Sulfolobus solfataricus

<400> 34
Tyr Leu Val Ile Lys Gly Tyr Tyr Ser Glu Leu Phe Thr Ser Ile Phe
  1               5                  10                  15

Ile Thr Asn Ser Phe Val Asp Phe Ile Phe Asn Phe Ile Ser Leu Tyr
             20                  25                  30

Val Ile Tyr Leu Ile Phe Gly Ser Arg Ala Gly Lys His Glu Tyr Gly
          35                  40                  45

Ile Phe Ile Leu Ala Gly Ile Leu Gly Asn Leu Leu Thr Val Ile Phe
      50                  55                  60

Tyr Ser Pro Phe Thr Leu Ser Ser Gly Ala Ser Gly Gly Ile Phe Gly
 65                  70                  75                  80

Leu Leu Ser Tyr Tyr Thr Phe Tyr Asp Phe Leu Lys Lys Asp Asn Leu
                 85                  90                  95

Gly Val Tyr Gly Leu Val Phe Leu Val Ser Val Phe Gly Val Ser Asp
            100                 105                 110

Leu Ile Phe Pro Asn Val Asn Val Val Ala His Ile Gly Gly Ile Leu
        115                 120                 125

Gly Gly Ile Met Tyr Ala Val Val Tyr Tyr Leu Ile Arg Ser
    130                 135                 140
```

<210> 35
<211> 156
<212> PRT
<213> Arabidopsis thaliana

<400> 35
Ile Phe Lys His Lys Asp Leu Lys Arg Leu Phe Leu Ser Ala Phe Tyr
1 5 10 15

His Val Asn Glu Pro His Leu Val Tyr Asn Met Met Ser Leu Leu Trp
20 25 30

Lys Gly Ile Lys Leu Glu Thr Ser Met Gly Ser Ser Glu Phe Ala Ser
35 40 45

Met Val Phe Thr Leu Ile Gly Met Ser Gln Gly Val Thr Leu Leu Leu
50 55 60

Ala Lys Ser Leu Leu Leu Leu Phe Asp Tyr Asp Arg Ala Tyr Tyr Asn
65 70 75 80

Glu Tyr Ala Val Gly Phe Ser Gly Val Leu Phe Ala Met Lys Val Val
85 90 95

Leu Asn Ser Gln Ala Glu Asp Tyr Ser Ser Val Tyr Gly Ile Leu Val
100 105 110

Pro Thr Lys Tyr Ala Ala Trp Ala Glu Leu Ile Leu Val Gln Met Phe
115 120 125

Val Pro Asn Ala Ser Phe Leu Gly His Leu Gly Gly Ile Leu Ala Gly
130 135 140

Ile Ile Tyr Leu Lys Leu Lys Gly Ser Tyr Ser Gly
145 150 155

<210> 36
<211> 10
<212> DNA
<213> Homo sapiens

<400> 36
tggccaataa 10

<210> 37
<211> 854
<212> PRT
<213> Homo sapiens

<400> 37
Met Ser Glu Ala Arg Arg Asp Ser Thr Ser Ser Leu Gln Arg Lys Lys
1 5 10 15

Pro Pro Trp Leu Lys Leu Asp Ile Pro Ser Ala Val Pro Leu Thr Ala
20 25 30

```
Glu Glu Pro Ser Phe Leu Gln Pro Leu Arg Arg Gln Ala Phe Leu Arg
         35                  40                  45

Ser Val Ser Met Pro Ala Glu Thr Ala His Ile Ser Ser Pro His His
     50                  55                  60

Glu Leu Arg Arg Pro Val Leu Gln Arg Gln Thr Ser Ile Thr Gln Thr
 65                  70                  75                  80

Ile Arg Arg Gly Thr Ala Asp Trp Phe Gly Val Ser Lys Asp Ser Asp
                 85                  90                  95

Ser Thr Gln Lys Trp Gln Arg Lys Ser Ile Arg His Cys Ser Gln Arg
            100                 105                 110

Tyr Gly Lys Leu Lys Pro Gln Val Leu Arg Glu Leu Asp Leu Pro Ser
        115                 120                 125

Gln Asp Asn Val Ser Leu Thr Ser Thr Glu Thr Pro Pro Pro Leu Tyr
    130                 135                 140

Val Gly Pro Cys Gln Leu Gly Met Gln Lys Ile Ile Asp Pro Leu Ala
145                 150                 155                 160

Arg Gly Arg Ala Phe Arg Val Ala Asp Asp Thr Ala Glu Gly Leu Ser
                165                 170                 175

Ala Pro His Thr Pro Val Thr Pro Gly Ala Ala Ser Leu Cys Ser Phe
            180                 185                 190

Ser Ser Ser Arg Ser Gly Phe His Arg Leu Pro Arg Arg Arg Lys Arg
        195                 200                 205

Glu Ser Val Ala Lys Met Ser Phe Arg Ala Ala Ala Ala Leu Met Lys
    210                 215                 220

Gly Arg Ser Val Arg Asp Gly Thr Phe Arg Arg Ala Arg Arg Ser Phe
225                 230                 235                 240

Thr Pro Ala Ser Phe Leu Glu Glu Asp Thr Thr Asp Phe Pro Asp Glu
                245                 250                 255

Leu Asp Thr Ser Phe Phe Ala Arg Glu Gly Ile Leu His Glu Glu Leu
            260                 265                 270

Ser Thr Tyr Pro Asp Glu Val Phe Glu Ser Pro Ser Glu Ala Ala Leu
        275                 280                 285

Lys Asp Trp Glu Lys Ala Pro Glu Gln Ala Asp Leu Thr Gly Gly Ala
    290                 295                 300

Leu Asp Arg Ser Glu Leu Glu Arg Ser His Leu Met Leu Pro Leu Glu
305                 310                 315                 320

Arg Gly Trp Arg Lys Gln Lys Glu Gly Ala Ala Ala Pro Gln Pro Lys
                325                 330                 335
```

```
Val Arg Leu Arg Gln Glu Val Val Ser Thr Ala Gly Pro Arg Arg Gly
            340                 345                 350

Gln Arg Ile Ala Val Pro Val Arg Lys Leu Phe Ala Arg Glu Lys Arg
        355                 360                 365

Pro Tyr Gly Leu Gly Met Val Gly Arg Leu Thr Asn Arg Thr Tyr Arg
    370                 375                 380

Lys Arg Ile Asp Ser Phe Val Lys Arg Gln Ile Glu Asp Met Asp Asp
385                 390                 395                 400

His Arg Pro Phe Phe Thr Tyr Trp Leu Thr Phe Val His Ser Leu Val
                405                 410                 415

Thr Ile Leu Ala Val Cys Ile Tyr Gly Ile Ala Pro Val Gly Phe Ser
            420                 425                 430

Gln His Glu Thr Val Asp Ser Val Leu Arg Asn Arg Gly Val Tyr Glu
        435                 440                 445

Asn Val Lys Tyr Val Gln Gln Glu Asn Phe Trp Ile Gly Pro Ser Ser
    450                 455                 460

Glu Ala Leu Ile His Leu Gly Ala Lys Phe Ser Pro Cys Met Arg Gln
465                 470                 475                 480

Asp Pro Gln Val His Ser Phe Ile Arg Ser Ala Arg Glu Arg Glu Lys
                485                 490                 495

His Ser Ala Cys Cys Val Arg Asn Asp Arg Ser Gly Cys Val Gln Thr
            500                 505                 510

Ser Glu Glu Glu Cys Ser Ser Thr Leu Ala Val Trp Val Lys Trp Pro
        515                 520                 525

Ile His Pro Ser Ala Pro Glu Leu Ala Gly His Lys Arg Gln Phe Gly
    530                 535                 540

Ser Val Cys His Gln Asp Pro Arg Val Cys Asp Glu Pro Ser Ser Glu
545                 550                 555                 560

Asp Pro His Glu Trp Pro Glu Asp Ile Thr Lys Trp Pro Ile Cys Thr
                565                 570                 575

Lys Asn Ser Ala Gly Asn His Thr Asn His Pro His Met Asp Cys Val
            580                 585                 590

Ile Thr Gly Arg Pro Cys Cys Ile Gly Thr Lys Gly Arg Cys Glu Ile
        595                 600                 605

Thr Ser Arg Glu Tyr Cys Asp Phe Met Arg Gly Tyr Phe His Glu Glu
    610                 615                 620

Ala Thr Leu Cys Ser Gln Val His Cys Met Asp Asp Val Cys Gly Leu
625                 630                 635                 640
```

```
Leu Pro Phe Leu Asn Pro Glu Val Pro Asp Gln Phe Tyr Arg Leu Trp
                645                 650                 655

Leu Ser Leu Phe Leu His Ala Gly Ile Leu His Cys Leu Val Ser Ile
            660                 665                 670

Cys Phe Gln Met Thr Val Leu Arg Asp Leu Glu Lys Leu Ala Gly Trp
        675                 680                 685

His Arg Ile Ala Ile Ile Tyr Leu Leu Ser Gly Val Thr Gly Asn Leu
    690                 695                 700

Ala Ser Ala Ile Phe Leu Pro Tyr Arg Ala Glu Val Gly Pro Ala Gly
705                 710                 715                 720

Ser Gln Phe Gly Ile Leu Ala Cys Leu Phe Val Glu Leu Phe Gln Ser
                725                 730                 735

Trp Gln Ile Leu Ala Arg Pro Trp Arg Ala Phe Phe Lys Leu Leu Ala
            740                 745                 750

Val Val Leu Phe Leu Phe Thr Phe Gly Leu Leu Pro Trp Ile Asp Asn
        755                 760                 765

Phe Ala His Ile Ser Gly Phe Ile Ser Gly Leu Phe Leu Ser Phe Ala
    770                 775                 780

Phe Leu Pro Tyr Ile Ser Phe Gly Lys Phe Asp Leu Tyr Arg Lys Arg
785                 790                 795                 800

Cys Gln Ile Ile Ile Phe Gln Val Val Phe Leu Gly Leu Leu Ala Gly
                805                 810                 815

Leu Val Val Leu Phe Tyr Val Tyr Pro Val Arg Cys Glu Trp Cys Glu
            820                 825                 830

Phe Leu Thr Cys Ile Pro Phe Thr Asp Lys Phe Cys Glu Lys Tyr Glu
        835                 840                 845

Leu Asp Ala Gln Leu His
    850
```

<210> 38
<211> 292
<212> PRT
<213> Homo sapiens

<400> 38

```
Met Asn Leu Asn Met Gly Arg Glu Met Lys Glu Glu Leu Glu Glu Glu
  1               5                  10                  15

Glu Lys Met Arg Glu Asp Gly Gly Gly Lys Asp Arg Ala Lys Ser Lys
             20                  25                  30

Lys Val His Arg Ile Val Ser Lys Trp Met Leu Pro Glu Lys Ser Arg
         35                  40                  45
```

```
Gly Thr Tyr Leu Glu Arg Ala Asn Cys Phe Pro Pro Pro Val Phe Ile
     50                  55                  60

Ile Ser Ile Ser Leu Ala Glu Leu Ala Val Phe Ile Tyr Tyr Ala Val
 65                  70                  75                  80

Trp Lys Pro Gln Lys Gln Trp Ile Thr Leu Asp Thr Gly Ile Leu Glu
                85                  90                  95

Ser Pro Phe Ile Tyr Ser Pro Glu Lys Arg Glu Glu Ala Trp Arg Phe
            100                 105                 110

Ile Ser Tyr Met Leu Val His Ala Gly Val Gln His Ile Leu Gly Asn
        115                 120                 125

Leu Cys Met Gln Leu Val Leu Gly Ile Pro Leu Glu Met Val His Lys
    130                 135                 140

Gly Leu Arg Val Gly Leu Val Tyr Leu Ala Gly Val Ile Ala Gly Ser
145                 150                 155                 160

Leu Ala Ser Ser Ile Phe Asp Pro Leu Arg Tyr Leu Val Gly Ala Ser
                165                 170                 175

Gly Gly Val Tyr Ala Leu Met Gly Gly Tyr Phe Met Asn Val Leu Val
            180                 185                 190

Asn Phe Gln Glu Met Ile Pro Ala Phe Gly Ile Phe Arg Leu Leu Ile
        195                 200                 205

Ile Ile Leu Ile Ile Val Leu Asp Met Gly Phe Ala Leu Tyr Arg Arg
    210                 215                 220

Phe Phe Val Pro Glu Asp Gly Ser Pro Val Ser Phe Ala Ala His Ile
225                 230                 235                 240

Ala Gly Gly Phe Ala Gly Met Ser Ile Gly Tyr Thr Val Phe Ser Cys
                245                 250                 255

Phe Asp Lys Ala Leu Leu Lys Asp Pro Arg Phe Trp Ile Ala Ile Ala
            260                 265                 270

Ala Tyr Leu Ala Cys Val Leu Phe Ala Val Phe Phe Asn Ile Phe Leu
        275                 280                 285

Ser Pro Ala Asn
    290
```

<210> 39
<211> 619
<212> PRT
<213> Homo sapiens

<400> 39

```
Met Ser Val Ala His Met Ser Leu Gln Ala Ala Ala Ala Leu Leu Lys
  1               5                  10                  15
```

```
Gly Arg Ser Val Leu Asp Ala Thr Gly Gln Arg Cys Arg Val Val Lys
            20                  25                  30

Arg Ser Phe Ala Phe Pro Ser Phe Leu Glu Glu Asp Val Val Asp Gly
        35                  40                  45

Ala Asp Thr Phe Asp Ser Ser Phe Phe Ser Lys Glu Glu Met Ser Ser
    50                  55                  60

Met Pro Asp Asp Val Phe Glu Ser Pro Pro Leu Ser Ala Ser Tyr Phe
65                  70                  75                  80

Arg Gly Ile Pro His Ser Ala Ser Pro Val Ser Pro Asp Gly Val Gln
                85                  90                  95

Ile Pro Leu Lys Glu Tyr Gly Arg Ala Pro Val Pro Gly Pro Arg Arg
            100                 105                 110

Gly Lys Arg Ile Ala Ser Lys Val Lys His Phe Ala Phe Asp Arg Lys
        115                 120                 125

Lys Arg His Tyr Gly Leu Gly Val Val Gly Asn Trp Leu Asn Arg Ser
    130                 135                 140

Tyr Arg Arg Ser Ile Ser Ser Thr Val Gln Arg Gln Leu Glu Ser Phe
145                 150                 155                 160

Asp Ser His Arg Pro Tyr Phe Thr Tyr Trp Leu Thr Phe Val His Val
                165                 170                 175

Ile Ile Thr Leu Leu Val Ile Cys Thr Tyr Gly Ile Ala Pro Val Gly
            180                 185                 190

Phe Ala Gln His Val Thr Thr Gln Leu Val Leu Arg Asn Lys Gly Val
        195                 200                 205

Tyr Glu Ser Val Lys Tyr Ile Gln Gln Glu Asn Phe Trp Val Gly Pro
    210                 215                 220

Ser Ser Ile Asp Leu Ile His Leu Gly Ala Lys Phe Ser Pro Cys Ile
225                 230                 235                 240

Arg Lys Asp Gly Gln Ile Glu Gln Leu Val Leu Arg Glu Arg Asp Leu
                245                 250                 255

Glu Arg Asp Ser Gly Cys Cys Val Gln Asn Asp His Ser Gly Cys Ile
            260                 265                 270

Gln Thr Gln Arg Lys Asp Cys Ser Glu Thr Leu Ala Thr Phe Val Lys
        275                 280                 285

Trp Gln Asp Asp Thr Gly Pro Pro Met Asp Lys Ser Asp Leu Gly Gln
    290                 295                 300

Lys Arg Thr Ser Gly Ala Val Cys His Gln Asp Pro Arg Thr Cys Glu
305                 310                 315                 320
```

```
Glu Pro Ala Ser Ser Gly Ala His Ile Trp Pro Asp Asp Ile Thr Lys
                325                 330                 335

Trp Pro Ile Cys Thr Glu Gln Ala Arg Ser Asn His Thr Gly Phe Leu
            340                 345                 350

His Met Asp Cys Glu Ile Lys Gly Arg Pro Cys Cys Ile Gly Thr Lys
        355                 360                 365

Gly Ser Cys Glu Ile Thr Thr Arg Glu Tyr Cys Glu Phe Met His Gly
    370                 375                 380

Tyr Phe His Glu Glu Ala Thr Leu Cys Ser Gln Val His Cys Leu Asp
385                 390                 395                 400

Lys Val Cys Gly Leu Leu Pro Phe Leu Asn Pro Glu Val Pro Asp Gln
                405                 410                 415

Phe Tyr Arg Leu Trp Leu Ser Leu Phe Leu His Ala Gly Val Val His
            420                 425                 430

Cys Leu Val Ser Val Val Phe Gln Met Thr Ile Leu Arg Asp Leu Glu
        435                 440                 445

Lys Leu Ala Gly Trp His Arg Ile Ala Ile Ile Phe Ile Leu Ser Gly
    450                 455                 460

Ile Thr Gly Asn Leu Ala Ser Ala Ile Phe Leu Pro Tyr Arg Ala Glu
465                 470                 475                 480

Val Gly Pro Ala Gly Ser Gln Phe Gly Leu Leu Ala Cys Leu Phe Val
                485                 490                 495

Glu Leu Phe Gln Ser Trp Pro Leu Leu Glu Arg Pro Trp Lys Ala Phe
            500                 505                 510

Leu Asn Leu Ser Ala Ile Val Leu Phe Leu Phe Ile Cys Gly Leu Leu
        515                 520                 525

Pro Trp Ile Asp Asn Ile Ala His Ile Phe Gly Phe Leu Ser Gly Leu
    530                 535                 540

Leu Leu Ala Phe Ala Phe Leu Pro Tyr Ile Thr Phe Gly Thr Ser Asp
545                 550                 555                 560

Lys Tyr Arg Lys Arg Ala Leu Ile Leu Val Ser Leu Leu Ala Phe Ala
                565                 570                 575

Gly Leu Phe Ala Ala Leu Val Leu Trp Leu Tyr Ile Tyr Pro Ile Asn
            580                 585                 590

Trp Pro Trp Ile Glu His Leu Thr Cys Phe Pro Phe Thr Ser Arg Phe
        595                 600                 605

Cys Glu Lys Tyr Glu Leu Asp Gln Val Leu His
    610                 615
```

<210> 40
<211> 404
<212> PRT
<213> Homo sapiens

<400> 40

```
Met Gly Glu His Pro Ser Pro Gly Pro Ala Val Ala Ala Cys Ala Glu
  1               5                  10                  15

Ala Glu Arg Ile Glu Glu Leu Glu Pro Glu Ala Glu Glu Arg Leu Pro
             20                  25                  30

Ala Ala Pro Glu Asp His Trp Lys Val Leu Phe Asp Gln Phe Asp Pro
         35                  40                  45

Gly Asn Thr Gly Tyr Ile Ser Thr Gly Lys Phe Arg Ser Leu Leu Glu
     50                  55                  60

Ser His Ser Ser Lys Leu Asp Pro His Lys Arg Glu Val Leu Leu Ala
 65                  70                  75                  80

Leu Ala Asp Ser His Ala Asp Gly Gln Ile Gly Tyr Gln Asp Phe Val
                 85                  90                  95

Ser Leu Met Ser Asn Lys Arg Ser Asn Ser Phe Arg Gln Ala Ile Leu
            100                 105                 110

Gln Gly Asn Arg Arg Leu Ser Ser Lys Ala Leu Leu Glu Glu Lys Gly
        115                 120                 125

Leu Ser Leu Ser Gln Arg Leu Ile Arg His Val Ala Tyr Glu Thr Leu
    130                 135                 140

Pro Arg Glu Ile Asp Arg Lys Trp Tyr Tyr Asp Ser Tyr Thr Cys Cys
145                 150                 155                 160

Pro Pro Pro Trp Phe Met Ile Thr Val Thr Leu Leu Glu Val Ala Phe
                165                 170                 175

Phe Leu Tyr Asn Gly Val Ser Leu Gly Gln Phe Val Leu Gln Val Thr
            180                 185                 190

His Pro Arg Tyr Leu Lys Asn Ser Leu Val Tyr His Pro Gln Leu Arg
        195                 200                 205

Ala Gln Val Trp Arg Tyr Leu Thr Tyr Ile Phe Met His Ala Gly Ile
    210                 215                 220

Glu His Leu Gly Leu Asn Val Val Leu Gln Leu Leu Val Gly Val Pro
225                 230                 235                 240

Leu Glu Met Val His Gly Ala Thr Arg Ile Gly Leu Val Tyr Val Ala
                245                 250                 255

Gly Val Val Ala Gly Ser Leu Ala Val Ser Val Ala Asp Met Thr Ala
            260                 265                 270
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Pro Val Val Gly Ser Ser Gly Gly Val Tyr Ala Leu Val Ser Ala His
        275                 280                 285

Leu Ala Asn Ile Val Met Asn Trp Ser Gly Met Lys Cys Gln Phe Lys
    290                 295                 300

Leu Leu Arg Met Ala Val Ala Leu Ile Cys Met Ser Met Glu Phe Gly
305                 310                 315                 320

Arg Ala Val Trp Leu Arg Phe His Pro Ser Ala Tyr Pro Pro Cys Pro
                325                 330                 335

His Pro Ser Phe Val Ala His Leu Gly Gly Val Ala Val Gly Ile Thr
            340                 345                 350

Leu Gly Val Val Val Leu Arg Asn Tyr Glu Gln Arg Leu Gln Asp Gln
        355                 360                 365

Ser Leu Trp Trp Ile Phe Val Ala Met Tyr Thr Val Phe Val Leu Phe
    370                 375                 380

Ala Val Phe Trp Asn Ile Phe Ala Tyr Thr Leu Leu Asp Leu Lys Leu
385                 390                 395                 400

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<210> 41
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<212> PRT
<213> Homo sapiens

<400> 41

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Trp Gly Ala Ser Val Gly Gly Arg Ser Cys Glu Glu Leu Thr Ala Val
             20                  25                  30

Leu Thr Pro Pro Gln Leu Leu Gly Arg Arg Phe Asn Phe Phe Ile Gln
         35                  40                  45

Gln Lys Cys Gly Phe Arg Lys Ala Pro Arg Lys Val Glu Pro Arg Arg
     50                  55                  60

Ser Asp Pro Gly Thr Ser Gly Glu Ala Tyr Lys Arg Ser Ala Leu Ile
 65                  70                  75                  80

Pro Pro Val Glu Glu Thr Val Phe Tyr Pro Ser Pro Tyr Pro Ile Arg
                 85                  90                  95

Ser Leu Ile Lys Pro Leu Phe Phe Thr Val Gly Phe Thr Gly Cys Ala
            100                 105                 110

Phe Gly Ser Ala Ala Ile Trp Gln Tyr Glu Ser Leu Lys Ser Arg Val
        115                 120                 125
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Gln Ser Tyr Phe Asp Gly Ile Lys Ala Asp Trp Leu Asp Ser Ile Arg
    130                 135                 140

Pro Gln Lys Glu Gly Asp Phe Arg Lys Glu Ile Asn Lys Trp Trp Asn
145                 150                 155                 160

Asn Leu Ser Asp Gly Gln Arg Thr Val Thr Gly Ile Ile Ala Ala Asn
                165                 170                 175

Val Leu Val Phe Cys Leu Trp Arg Val Pro Ser Leu Gln Arg Thr Met
            180                 185                 190

Ile Arg Tyr Phe Thr Ser Asn Pro Ala Ser Lys Val Leu Cys Ser Pro
        195                 200                 205

Met Leu Leu Ser Thr Phe Ser His Phe Ser Leu Phe His Met Ala Ala
    210                 215                 220

Asn Met Tyr Val Leu Trp Ser Phe Ser Ser Ser Ile Val Asn Ile Leu
225                 230                 235                 240

Gly Gln Glu Gln Phe Met Ala Val Tyr Leu Ser Ala Gly Val Ile Ser
                245                 250                 255

Asn Phe Val Ser Tyr Leu Gly Lys Val Ala Thr Gly Arg Tyr Gly Pro
            260                 265                 270

Ser Leu Gly Ala Ser Gly Ala Ile Met Thr Val Leu Ala Ala Val Cys
        275                 280                 285

Thr Lys Ile Pro Glu Gly Arg Leu Ala Ile Ile Phe Leu Pro Met Phe
    290                 295                 300

Thr Phe Thr Ala Gly Asn Ala Leu Lys Ala Ile Ile Ala Met Asp Thr
305                 310                 315                 320

Ala Gly Met Ile Leu Gly Trp Lys Phe Phe Asp His Ala Ala His Leu
                325                 330                 335

Gly Gly Ala Leu Phe Gly Ile Trp Tyr Val Thr Tyr Gly His Glu Leu
            340                 345                 350

Ile Trp Lys Asn Arg Glu Pro Leu Val Lys Ile Trp His Glu Ile Arg
        355                 360                 365

Thr Asn Gly Pro Lys Lys Gly Gly Gly Ser Lys
    370                 375


<210> 42
<211> 315
<212> PRT
<213> Homo sapiens

<400> 42
Met Gln Arg Arg Ser Arg Gly Ile Asn Thr Gly Leu Ile Leu Leu Leu
  1               5                  10                  15
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Ser Gln Ile Phe His Val Gly Ile Asn Asn Ile Pro Pro Val Thr Leu
            20                  25                  30

Ala Thr Leu Ala Leu Asn Ile Trp Phe Phe Leu Asn Pro Gln Lys Pro
        35                  40                  45

Leu Tyr Ser Ser Cys Leu Ser Val Glu Lys Cys Tyr Gln Gln Lys Asp
    50                  55                  60

Trp Gln Arg Leu Leu Leu Ser Pro Leu His His Ala Asp Asp Trp His
65                  70                  75                  80

Leu Tyr Phe Asn Met Ala Ser Met Leu Trp Lys Gly Ile Asn Leu Glu
                85                  90                  95

Arg Arg Leu Gly Ser Arg Trp Phe Ala Tyr Val Ile Thr Ala Phe Ser
            100                 105                 110

Val Leu Thr Gly Val Val Tyr Leu Leu Leu Gln Phe Ala Val Ala Glu
        115                 120                 125

Phe Met Asp Glu Pro Asp Phe Lys Arg Ser Cys Ala Val Gly Phe Ser
    130                 135                 140

Gly Val Leu Phe Ala Leu Lys Val Leu Asn Asn His Tyr Cys Pro Gly
145                 150                 155                 160

Gly Phe Val Asn Ile Leu Gly Phe Pro Val Pro Asn Arg Phe Ala Cys
                165                 170                 175

Trp Val Glu Leu Val Ala Ile His Leu Phe Ser Pro Gly Thr Ser Phe
            180                 185                 190

Ala Gly His Leu Ala Gly Ile Leu Val Gly Leu Met Tyr Thr Gln Gly
        195                 200                 205

Pro Leu Lys Lys Ile Met Glu Ala Cys Ala Gly Gly Phe Ser Ser Ser
    210                 215                 220

Val Gly Tyr Pro Gly Arg Gln Tyr Tyr Phe Asn Ser Ser Gly Ser Ser
225                 230                 235                 240

Gly Tyr Gln Asp Tyr Tyr Pro His Gly Arg Pro Asp His Tyr Glu Glu
                245                 250                 255

Ala Pro Arg Asn Tyr Asp Thr Tyr Thr Ala Gly Leu Ser Glu Glu Glu
            260                 265                 270

Gln Leu Glu Arg Ala Leu Gln Ala Ser Leu Trp Asp Arg Gly Asn Thr
        275                 280                 285

Arg Asn Ser Pro Pro Pro Tyr Gly Phe His Leu Ser Pro Glu Glu Met
    290                 295                 300

Arg Arg Gln Arg Leu His Arg Phe Asp Ser Gln
305                 310                 315
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<210> 43
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 43
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<210> 44
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 44
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<210> 45
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 45
Lys Met Ser Phe Arg Ala Ala Ala Ala
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<210> 46
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
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<400> 46
Leu Thr Ala Glu Glu Pro Ser Phe Leu
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<210> 47
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 47
Ser Gln His Glu Thr Val Asp Ser Val
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<210> 48
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<212> PRT
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Gly Val Tyr Glu Asn Val Lys Tyr Val
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<211> 9
<212> PRT
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Tyr Val Gln Gln Glu Asn Phe Trp Ile
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<211> 9
<212> PRT
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<210> 51
<211> 30
<212> PRT
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<400> 51
Arg Gly Arg Ala Phe Arg Val Ala Asp Asp Thr Ala Glu Gly Leu Ser
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Ala Pro His Thr Pro Val Thr Pro Gly Ala Ala Ser Leu Cys
20 25 30

<210> 52
<211> 30
<212> PRT
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic peptide

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Val Lys Tyr Val Gln Gln Glu Asn Phe Trp Ile Gly Pro Ser Ser Glu
1 5 10 15

Ala Leu Ile His Leu Gly Ala Lys Phe Ser Pro Cys Met Arg
20 25 30

<210> 53
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 53
Pro Val Arg Cys Glu Trp Cys Glu Phe Leu Thr Cys Ile Pro Phe Thr
1 5 10 15

Asp Lys Phe Cys Glu Lys Tyr Glu Leu Asp Ala Gln Leu His
20 25 30

<210> 54
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 54
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<210> 55
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 55
ccctgggatc ctggtggcag acagag 26

<210> 56
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 56
ccggcgtcga ctcagtggag ctgagcgtc 29

<210> 57
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 57
caccaggatc ccagggtgtg tgatga 26

<210> 58
<211> 6559
<212> DNA
<213> Homo sapiens

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gtcattagtt catagcccat atatggagtt ccgcgttaca taacttacgg taaatggccc 240
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<211> 29
<212> DNA
<213> Artificial Sequence

<220>
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<210> 60
<211> 5653
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<213> Homo sapiens

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<210> 63
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

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<210> 64
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<213> Homo sapiens

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<210> 65
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

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<210> 66
<211> 6185
<212> DNA
<213> Homo sapiens

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<210> 67
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic primer

<400> 67
cgcggcatcg atgtccatgt cctcgatctg 30

<210> 68
<211> 4820
<212> DNA
<213> Homo sapiens

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<211> 4736
<212> DNA
<213> Homo sapiens

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<400> 70
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<210> 71

<400> 71
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<210> 72
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 72
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<210> 73
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic oligonucleotide

<400> 73
aactgttgag gagcccatgg a 21

<210> 74

<400> 74
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<210> 75

<400> 75
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<210> 76
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 76
Ala Leu Trp Val Leu Gly Leu Cys Cys
1 5

<210> 77
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 77
Val Leu Gly Leu Cys Cys Val Leu Leu
1 5

<210> 78
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 78
Leu Leu His Val Thr Asp Thr Gly Val
1 5

<210> 79
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 79
Ser Glu Leu Ile Gly Gln Phe Gly Val
1 5

<210> 80
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 80
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Lys Ser Arg Glu Gly Ser Arg Thr Asp Asp Glu Val Val Gln
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<210> 81
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 81
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Asn Asp Thr Gln His Ile Trp Glu Ser Asp Ser Asn Glu Phe
20 25 30

<210> 82
<211> 30
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 82
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1 5 10 15

Pro Leu Leu Asn Trp Met Lys Asp Lys Ala Leu Lys Asp Lys
20 25 30

<210> 83
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 83
Met Met Pro Lys Tyr Leu Asn Phe Val
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<210> 84
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 84
Lys Leu Tyr Val Arg Arg Val Phe Ile
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<210> 85
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 85
Arg Leu Leu Lys Lys Gly Tyr Glu Val
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<210> 86
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 86
Phe Leu Val Ala Asp Lys Val Ile Val
1 5

<210> 87
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 87
Leu Leu His Val Thr Asp Thr Gly Val
1 5

<210> 88
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 88
Lys Glu Ala Glu Ser Ser Pro Phe Val
1 5

<210> 89
<211> 9
<212> PRT
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<220>
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<400> 89
Arg Leu Thr Glu Ser Pro Cys Ala Leu
1 5

<210> 90
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
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<400> 90
Val Thr Phe Lys Ser Ile Leu Phe Val
1 5

<210> 91
<211> 9
<212> PRT
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<220>
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<400> 91
Ala Leu Trp Val Leu Gly Leu Cys Cys
1 5

<210> 92
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 92
Val Leu Gly Leu Cys Cys Val Leu Leu
1 5

<210> 93
<211> 9
<212> PRT
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<220>
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<400> 93
Ser Glu Leu Ile Gly Gln Phe Gly Val
1 5

<210> 94
<211> 9
<212> PRT
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<220>
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<400> 94
Arg Met Leu Arg Leu Ser Leu Asn Ile
1 5

<210> 95
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 95
Leu Gln Gln His Lys Leu Leu Lys Val
1 5

<210> 96
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 96
Tyr Val Trp Ser Ser Lys Thr Glu Thr
1 5

<210> 97
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 97
Leu Glu Leu Asp Thr Ile Lys Asn Leu
1 5

<210> 98
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 98
Phe Ile Thr Asp Asp Phe His Asp Met
1 5

<210> 99
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 99
Lys Thr Leu Asp Met Ile Lys Lys Ile
1 5

<210> 100
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 100
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1 5

<210> 101
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 101
Tyr Leu Asn Phe Val Lys Gly Val Val
1 5

<210> 102
<211> 9
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 102
Val Gly Phe Tyr Ser Ala Phe Leu Val
1 5

<210> 103
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<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 103
Lys Asp Glu Leu
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<210> 104
<211> 5
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide motif

<400> 104
Gly Ala Ser Gly Gly
1 5

<210> 105
<211> 5
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide motif

<400> 105
Gly Asp Ser Gly Gly
1 5

<210> 106
<211> 23
<212> PRT
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<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 106
Trp Leu Thr Phe Val His Ser Leu Val Thr Ile Leu Ala Val Cys Ile
1 5 10 15

Tyr Gly Ile Ala Pro Val Gly
20

<210> 107
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 107
Leu Trp Leu Ser Leu Phe Leu His Ala Gly Ile Leu His Cys Leu Val
1 5 10 15

Ser Ile Cys Phe Gln Met Thr
20

<210> 108
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 108
Leu Ser Gly Val Thr Gly Asn Leu Ala Ser Ala Ile Phe Leu Pro Tyr
1 5 10 15

Arg Ala Glu Val Gly Pro Ala
20

<210> 109
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 109
Trp Arg Ala Phe Phe Lys Leu Leu Ala Val Val Leu Phe Leu Phe Thr
1 5 10 15

Phe Gly Leu Leu Pro Trp Ile
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<210> 110
<211> 22
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 110
Ile Ser Gly Phe Ile Ser Gly Leu Phe Leu Ser Phe Ala Phe Leu Pro
1 5 10 15

Tyr Ile Ser Phe Gly Lys
20

<210> 111
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic peptide

<400> 111
Gln Ile Ile Ile Phe Gln Val Val Phe Leu Gly Leu Leu Ala Gly Leu
1 5 10 15

Val Val Leu Phe Tyr Val Tyr
20